

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

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Application Serial Number: 10/562,601
Source: FWO
Date Processed by STIC: 3/5/07

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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Revised 01/10/06



IFWO

RAW SEQUENCE LISTING

DATE: 03/05/2007

PATENT APPLICATION: US/10/562,601

TIME: 13:44:39

Input Set : N:\efs\03_05_07\10562601_efs\sequencelisting.txt

Output Set: N:\CRF4\03052007\J562601.raw

3 <110> APPLICANT: Hengst den, Christiaan D.
 4 Gajic, Olivera
 5 Kuipers, Oscar P.
 6 Kok, Jan
 7 Sikkema, Jan
 8 Geurts, Johannes M.W.
 9 Nauta, Arjen
 11 <120> TITLE OF INVENTION: Methods and means for regulating gene expression
 13 <130> FILE REFERENCE: P63590US00
 15 <140> CURRENT APPLICATION NUMBER: US 10/562,601
 16 <141> CURRENT FILING DATE: 2005-12-28
 18 <150> PRIOR APPLICATION NUMBER: PCT/NL2004/000474
 19 <151> PRIOR FILING DATE: 2004-07-02
 21 <150> PRIOR APPLICATION NUMBER: EP 03077074.7
 22 <151> PRIOR FILING DATE: 2003-07-02
 24 <160> NUMBER OF SEQ ID NOS: 234
 27 <170> SOFTWARE: PatentIn Ver. 3.3

Does Not Comply
Corrected Diskette Needed

see p. 2, 4

ERRORED SEQUENCES

2299 <210> SEQ ID NO: 179
 2300 <211> LENGTH: 26
 2301 <212> TYPE: DNA
 2302 <213> ORGANISM: Artificial Sequence
 2304 <220> FEATURE:
 2305 <223> OTHER INFORMATION: presence of the putative CodY box in arcA
 E--> 2307 <400> SEQUENCE: 179 see p. 2
 2308 atatttttttg aaaaattttt taaaaa

26

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<210> 179
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> presence of the putative CodY box in arcA

Ⓔ <400> 179
atatttttttg aaaaattttt taaaaa

26

delete extra opening bracket

VERIFICATION SUMMARY

DATE: 03/05/2007

PATENT APPLICATION: US/10/562,601

TIME: 13:44:40

Input Set : N:\efs\03_05_07\10562601_efs\sequencelisting.txt

Output Set: N:\CRF4\03052007\J562601.raw

L:2307 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:179 differs:178
L:2739 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2732
L:2744 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2737
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2742
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2747
L:2914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:0
L:2931 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:228 after pos.:0
L:2960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:230 after pos.:0

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<210> 15
<211> 8
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<213> Artificial Sequence

<220>
<223> consensus sequence GTP binding motif in CodY homologs

<220>
<221> Variant
<222> (2)..(2)
<223> Arg

if the amino acid at this location can represent another amino acid, use "Xaa" instead, and

specify in <2207-22237 section

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<221> Variant
<222> (3)..(3)
<223> Met, Gly, Ile, Lys, Gln

same

<220>
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<222> (4)..(4)
<223> Gly

same

<220>
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<222> (5)..(5)
<223> Thr

same

<220>
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<222> (7)..(7)
<223> Ser

same

<400> 15
Gly Gly Glu Arg Leu Gly Thr Thr
1 5